

SEQUENCE LISTING

<110> Delcayre, Alain

<120> Compositions isolated from *M. vaccae* and
their use in modulation of immune responses.

<130> 11000.1047c2

<150> US 10/051,325

<151> 2002-01-18

<150> US 09/455,960

<151> 1999-12-06

<160> 31

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<212> DNA

<213> *Mycobacterium vaccae*

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taccccaaca	tcaggacgct	gctcggcaag	gcggtcgaga	tcgaccccg	ccgcccgggtg	240
gtgaccgcga	tgagaccgga	cgaatccacg	ttcacgctcg	actacgacgt	gctcgtcgtc	300
gccgcccgg	tcgacgagtc	ctatttcggc	aagcgtcatt	tcgcgaggag	ggcgccaggg	360
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<212> DNA

<213> *Mycobacterium vaccae*

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cttcgcgatg	ctgtggctca	acaaggccgt	cgccaccgac	gccgaccggg	tcgccaccgc	180
cgaacgggtg	gcgcgcatgc	tggtcacccc	gaaccggccc	gccgggaccc	ccgacgagta	240
gcgtcggcgt	catgactgac	gctgcgatca	ccgacatccc	gctcaccacc	ctggacggcc	300
ggcccaccac	gctcgcggag	ttggccgacg	gcgcccgcgt	gggtgtcaat	gtcgcctcca	360
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gcggactcac	cgtgatcggc	gtgccgtgca	accagtccat	ggggcaggag	cccgggaccg	480
ccgaggagat	ccagacgttc	tgctcgacga	cctacgacgt	gacgttccc	ctgctggaga	540
agaccgacgt	caacggggccc	ggcaggcatc	cgctctacgc	cgagctggcc	cgcgccaccg	600
acgaggacgg	cgaggccggc	gacgtgcagt	ggaacttcga	gaagtccctg	ctcgccccgg	660
gcggcaaagt	ggtcaggcgt	ttccgtcccc	gcaccgcccc	ggacgcccc	gaggtgatct	720
cggccatcga	agacgtcttg	ccccgatagc	cgaagcgaca	cctgggcgcg	cggtgtcgtc	780
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atctccggaa	tcagtgaccg	gacctcgggt	gaggtcgccc	agttccggag	tcgctcgac	900
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<212> DNA

<213> Mycobacterium vaccae

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gtgatcggcg	tgccgtgcaa	ccagtccatg	gggcaggagc	ccgggaccgc	cgaggagatc	240
cagacgttct	gctcgacgac	ctacgacgtg	acgttccccg	tgctggagaa	gaccgacgtc	300
aacggggccc	gcaggcatcc	gctctacgcc	gagctggccc	gcgccaccga	cgaggacggc	360
gaggccggcg	acgtgcagtg	gaacttcgag	aagtccctgc	tcgccccggg	cggcaaagtg	420
gtcaggcggt	tcggtccccg	caccgccccg	gacgcccccg	aggtgatctc	ggccatcgaa	480
gacgtcttgc	cccgatag					498

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cggctggacc	gggacgcggc	caccgtcgac	tggctgatcg	accggcgccg	gcgcggcgat	180
gccgtcgtgc	tgacgggttt	cgacgaggcc	cggaccacag	cgcgccgcgg	tgagttcgcg	240
acgtgccccg	cgcacgaggc	gaacctgcgc	ctgatggccg	ccgacaggat	catggagcac	300
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aaagtactgc	cccgcaatgg	gtttcgcgtc	cttgccgggg	tggcgggcat	cgtcgacctg	420
gtcgggtgggc	acactgtgcg	cgcgcgggtg	ctcggcatcg	gcgggggctt	cctcgccgaa	480

c c g t g g t g g t	g c c g g a c g c t	g g t g c t g g c c	g c c g a a c g c a	c g g c a c g c c g	c g g c g g a a c c	540
g t g c g g c t g a	c g g t g t c g g c	c c g g c a g c t g	a g c c g c c c g g	g t c c c c g g c a	a a c c c t g t c t c	600
g a c g c c g t c g	a a c t g g c g a t	g c t g c a c t c g	g g c g c g g c c a	c g g t c t a c c g	g t g g c a c c c c	660
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c g c g g c c g a a	g c g t g g t g a t	c g t c g a c c a g	g a g a a c g c g g	c c a a t g t c g g	c g g c c a g g c g	120
t t c t g g t c g t	t c g g c g g g c t	g t t c t t c g t c	g a c a g c c c g g	a g c a g c g g c g	c a t g g g c a t c	180
c g g g a c a g t c	a c g a g c t c g c	g c t g c a g g a c	t g g t c g g c t	c g g c c g g g t t	c g a c c g g c c c	240
g a g g a c c a c t	g g c c g c g g c t	g t g g g c c c a c	g c c t a c g t c g	a c t t c g c c g c	c g g c g a g a a g	300
c g c a g c t g g c	t g c g c g a g c g	c g g t c t g c a g	a c c t t c g c g c	t g g t c g g c t g	g g c c g a a c g c	360
g g c g g c t a c g	g g g c c a a c g g	g c a c g g c a a c	t c g g t g c c g c	g c t t c c a c a t	c a c g t g g g g c	420
a c c g g g c c c g	c g c t g g t c g a	c a t c t t c g c g	c g g c g g t t g a	c c g g g g t g c c	g c g g g t g c g g	480
t t c g t c c a c c	g g c a c c g g g t	g g a c g a g c t g	a t c g t c g a g g	a c g g t g c g g t	g g t c g g g g t g	540
c g c g g a g c c g	t a c t g g a a c c	g t c g t c g g c g	g t a c c c g g t g	c g g a a t c c t c	c c g c g a g g t c	600
g t c g g c g a c t	t c g a g a t g c g	g g c g c a g g c g	g t g a t c g t g g	c c a g c g g c g g	g a t c g g g g g c	660
a a c c a c g a c c	t g g t g c g c a a	a t a c t g g c c c	a a g c g g a t g g	g a c g g g t g c c	c g a a c a a c t g	720
c t c a g c g g t g	t g c c c g c g c a	c g t c g a c g g a	c g c a t g c t g c	a g a t c t c g g a	g a c c g c g g g t	780
g c c a g c g t c a	t c a a c a a a g a	c c g g a t g t g g	c a c t a c a c c g	a g g g c a t c a c	c a a c t a c g a c	840
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			20					25					30				
Phe	Gln	Pro	Leu	Leu	Tyr	Gln	Cys	Ala	Thr	Gly	Thr	Leu	Ser	Ile	Ala		
			35				40					45					
His	Ile	Ser	Arg	Pro	Leu	Arg	Glu	Glu	Phe	Ala	Arg	Tyr	Pro	Asn	Ile		
	50					55				60							
Arg	Thr	Leu	Leu	Gly	Lys	Ala	Val	Glu	Ile	Asp	Pro	Asp	Arg	Arg	Val		
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Val	Thr	Ala	Met	Arg	Pro	Asp	Glu	Ser	Thr	Phe	Thr	Leu	Asp	Tyr	Asp		
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Val	Leu	Val	Val	Ala	Ala	Gly	Met	Gln	Gln	Ser	Tyr	Phe	Gly	Lys	Arg		
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His	Phe	Ala	Glu	Trp	Ala	Pro	Gly	Met	Lys	Thr	Leu	Asp	Asp	Ala	Leu		
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Asn	Val	Ala	Ser	Lys	Cys	Gly	Leu	Thr	Pro	Gln	Tyr	Thr	Ala	Leu	Glu
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Gln	Leu	Ala	Lys	Asp	Tyr	Arg	Glu	Arg	Gly	Leu	Thr	Val	Ile	Gly	Val
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Gln	Thr	Phe	Cys	Ser	Thr	Thr	Tyr	Asp	Val	Thr	Phe	Pro	Leu	Leu	Glu
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Lys	Thr	Asp	Val	Asn	Gly	Pro	Gly	Arg	His	Pro	Leu	Tyr	Ala	Glu	Leu
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Ala	Arg	Ala	Thr	Asp	Glu	Asp	Gly	Glu	Ala	Gly	Asp	Val	Gln	Trp	Asn
		115					120					125			
Phe	Glu	Lys	Phe	Leu	Leu	Ala	Pro	Gly	Gly	Lys	Val	Val	Arg	Arg	Phe
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Asp	Val	Leu	Pro	Arg											
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<213> Mycobacterium vaccae

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			20					25					30		
Ala	Pro	Arg	Leu	Lys	Gly	Gly	Tyr	Arg	Leu	Asp	Arg	Asp	Ala	Ala	Thr
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Val	Asp	Trp	Leu	Ile	Asp	Arg	Arg	Arg	Arg	Gly	Asp	Ala	Val	Val	Leu
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His	Gly	Phe	Asp	Glu	Ala	Arg	Thr	Thr	Ala	Arg	Arg	Gly	Glu	Phe	Ala
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Thr	Leu	Pro	Ala	His	Glu	Ala	Asn	Leu	Arg	Leu	Met	Ala	Ala	Asp	Arg
			85						90					95	
Ile	Met	Glu	His	Leu	Asp	Leu	Arg	Thr	Arg	Ile	Phe	Ala	Ala	Pro	Gly
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Trp	Asn	Val	Ser	Pro	Gly	Ala	Leu	Lys	Val	Leu	Pro	Arg	Asn	Gly	Phe
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Thr	Val	Arg	Ala	Arg	Val	Leu	Gly	Ile	Gly	Gly	Gly	Phe	Leu	Ala	Glu
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Pro	Trp	Trp	Cys	Arg	Thr	Leu	Val	Leu	Ala	Ala	Glu	Arg	Thr	Ala	Arg
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			180					185					190		
Pro	Gly	Pro	Arg	Gln	Thr	Leu	Leu	Asp	Ala	Val	Glu	Leu	Ala	Met	Leu
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210
Thr Glu Ala Ala
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215

220

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<213> Mycobacterium vaccae

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Phe	Val	Asp	Ser	Pro	Glu	Gln	Arg	Arg	Met	Gly	Ile	Arg	Asp	Ser	His
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Glu	Asp	His	Trp	Pro	Arg	Leu	Trp	Ala	His	Ala	Tyr	Val	Asp	Phe	Ala
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Ala	Gly	Glu	Lys	Arg	Ser	Trp	Leu	Arg	Glu	Arg	Gly	Leu	Gln	Thr	Phe
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Ala	Leu	Val	Gly	Trp	Ala	Glu	Arg	Gly	Gly	Tyr	Gly	Ala	Asn	Gly	His
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Leu	Val	Asp	Ile	Phe	Ala	Arg	Arg	Leu	Thr	Gly	Val	Pro	Arg	Val	Arg
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Leu	Ser	Gly	Val	Pro	Ala	His	Val	Asp	Gly	Arg	Met	Leu	Gln	Ile	Ser
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